

#10



PCT09

RAW SEQUENCE LISTING

DATE: 06/24/2002

PATENT APPLICATION: US/09/890,811A

TIME: 17:40:08

Input Set : N:\Crf3\Refhold\I890811A.raw

Output Set: N:\CRF3\06242002\I890811A.raw

ENTERED

C--> 1 <110> APPLICANT: Zhu, Qun
2 Famodu, Omolayo O.
3 <120> TITLE OF INVENTION: SPFl-Related Transcription Factors
4 <130> FILE REFERENCE: BB1436 PCT
5 <140> CURRENT APPLICATION NUMBER: US/09/890,811A
6 <141> CURRENT FILING DATE: 2002-05-07
7 <150> PRIOR APPLICATION NUMBER: 60/174325
8 <151> PRIOR FILING DATE: 2000-01-04
9 <160> NUMBER OF SEQ ID NOS: 14
10 <170> SOFTWARE: Microsoft Office 97
12 <210> SEQ ID NO: 1
13 <211> LENGTH: 512
14 <212> TYPE: DNA
15 <213> ORGANISM: Zea mays
16 <220> FEATURE:
17 <221> NAME/KEY: unsure
18 <222> LOCATION: (368)
19 <223> OTHER INFORMATION: n=a,c,g or t
20 <220> FEATURE:
21 <221> NAME/KEY: unsure
22 <222> LOCATION: (387)
23 <223> OTHER INFORMATION: n=a,c,g or t
24 <220> FEATURE:
25 <221> NAME/KEY: unsure
26 <222> LOCATION: (405)
27 <223> OTHER INFORMATION: n=a,c,g or t
28 <220> FEATURE:
29 <221> NAME/KEY: unsure
30 <222> LOCATION: (408)
31 <223> OTHER INFORMATION: n=a,c,g or t
32 <220> FEATURE:
33 <221> NAME/KEY: unsure
34 <222> LOCATION: (445)
35 <223> OTHER INFORMATION: n=a,c,g or t
36 <220> FEATURE:
37 <221> NAME/KEY: unsure
38 <222> LOCATION: (447)
39 <223> OTHER INFORMATION: n=a,c,g or t
40 <220> FEATURE:
41 <221> NAME/KEY: unsure
42 <222> LOCATION: (453)
43 <223> OTHER INFORMATION: n=a,c,g or t
44 <220> FEATURE:

RAW SEQU LISTING D 06/24/2002
 PATENT APPLICATION: US/09/890,811A TIME: 17:40:08

Input Set : N:\Crf3\Refhold\I890811A.raw
 Output Set: N:\CRF3\06242002\I890811A.raw

45 <221> NAME/KEY: unsure
 46 <222> LOCATION: (462)
 47 <223> OTHER INFORMATION: n=a,c,g or t
 48 <220> FEATURE:
 49 <221> NAME/KEY: unsure
 50 <222> LOCATION: (489)..(491)
 51 <223> OTHER INFORMATION: n=a,c,g or t
 52 <220> FEATURE:
 53 <221> NAME/KEY: unsure
 54 <222> LOCATION: (500)
 55 <223> OTHER INFORMATION: n=a,c,g or t
 56 <400> SEQUENCE: 1
 57 ctataattgg cgcaaatatg gacagaagca tgtcaaggga agtgaaaatc ctagaagtta 60
 58 ttacaagtgc actcatccta attgtgaagt taaaaagcta ttagagcggt cgcttgatgg 120
 59 tcagattact gaagttgttt ataaagggca tcataatcat cccaagcccc aaccaaatag 180
 60 aaggttagct gctggtgcag ttccttcaag ccaggctgaa gaaagatacg atggtgtggc 240
 61 acctattgaa gacaaacctt caaatattta ttccaacctc tgtaaccaag cacattcagc 300
 62 tggcatgggt gataatgttc cgggtccagc aagtgatgat gatgttgatg ctggaggtgg 360
 63 aagaccnacc cctgggggga tgactcnaat gatgatgatg acttnggnct caaaaacgca 420
 64 aggaaaatgg aatctgccgg gtatncnatg ccnggctttg antggggtaa accaaatccc 480
 65 ggagccccnn nttccttttn aaaaactggt tt 512
 67 <210> SEQ ID NO: 2
 68 <211> LENGTH: 170
 69 <212> TYPE: PRT
 70 <213> ORGANISM: Zea mays
 71 <220> FEATURE:
 72 <221> NAME/KEY: unsure
 73 <222> LOCATION: (123)
 74 <223> OTHER INFORMATION: Xaa = ANY AMINO ACID
 75 <220> FEATURE:
 76 <221> NAME/KEY: unsure
 77 <222> LOCATION: (129)
 78 <223> OTHER INFORMATION: Xaa = ANY AMINO ACID
 79 <220> FEATURE:
 80 <221> NAME/KEY: unsure
 81 <222> LOCATION: (135)..(136)
 82 <223> OTHER INFORMATION: Xaa = ANY AMINO ACID
 83 <220> FEATURE:
 84 <221> NAME/KEY: unsure
 85 <222> LOCATION: (148)..(149)
 86 <223> OTHER INFORMATION: Xaa = ANY AMINO ACID
 87 <220> FEATURE:
 88 <221> NAME/KEY: unsure
 89 <222> LOCATION: (151)
 90 <223> OTHER INFORMATION: Xaa = ANY AMINO ACID
 91 <220> FEATURE:
 92 <221> NAME/KEY: unsure
 93 <222> LOCATION: (154)
 94 <223> OTHER INFORMATION: Xaa = ANY AMINO ACID

RAW SEQU LISTING D 06/24/2002
 PATENT APPLICATION: US/09/890,811A TIME: 17:40:08

Input Set : N:\Crf3\Refhold\I890811A.raw
 Output Set: N:\CRF3\06242002\I890811A.raw

```

95 <220> FEATURE:
96 <221> NAME/KEY: unsure
97 <222> LOCATION: (163)..(164)
98 <223> OTHER INFORMATION: Xaa = ANY AMINO ACID
99 <220> FEATURE:
100 <221> NAME/KEY: unsure
101 <222> LOCATION: (167)
102 <223> OTHER INFORMATION: Xaa = ANY AMINO ACID
103 <400> SEQUENCE: 2
104   Tyr Asn Trp Arg Lys Tyr Gly Gln Lys His Val Lys Gly Ser Glu Asn
105       1           5           10           15
106   Pro Arg Ser Tyr Tyr Lys Cys Thr His Pro Asn Cys Glu Val Lys Lys
107           20           25           30
108   Leu Leu Glu Arg Ser Leu Asp Gly Gln Ile Thr Glu Val Val Tyr Lys
109           35           40           45
110   Gly His His Asn His Pro Lys Pro Gln Pro Asn Arg Arg Leu Ala Ala
111       50           55           60
112   Gly Ala Val Pro Ser Ser Gln Ala Glu Glu Arg Tyr Asp Gly Val Ala
113       65           70           75           80
114   Pro Ile Glu Asp Lys Pro Ser Asn Ile Tyr Ser Asn Leu Cys Asn Gln
115           85           90           95
116   Ala His Ser Ala Gly Met Val Asp Asn Val Pro Gly Pro Ala Ser Asp
117           100          105          110
W--> 118   Asp Asp Val Asp Ala Gly Gly Gly Arg Pro Xaa Pro Gly Gly Met Thr
119           115          120          125
W--> 120   Xaa Met Met Met Met Thr Xaa Xaa Ser Lys Thr Gln Gly Lys Trp Asn
121           130          135          140
W--> 122   Leu Pro Gly Xaa Xaa Cys Xaa Ala Leu Xaa Gly Val Asn Gln Ile Pro
123       145          150          155          160
W--> 124   Glu Pro Xaa Xaa Pro Phe Xaa Lys Leu Phe
125           165          170
127 <210> SEQ ID NO: 3
128 <211> LENGTH: 717
129 <212> TYPE: DNA
130 <213> ORGANISM: Zea mays
131 <220> FEATURE:
132 <221> NAME/KEY: unsure
133 <222> LOCATION: (11)
134 <223> OTHER INFORMATION: n=a,c,g or t
135 <220> FEATURE:
136 <221> NAME/KEY: unsure
137 <222> LOCATION: (15)
138 <223> OTHER INFORMATION: n=a,c,g or t
139 <220> FEATURE:
140 <221> NAME/KEY: unsure
141 <222> LOCATION: (610)
142 <223> OTHER INFORMATION: n=a,c,g or t
143 <220> FEATURE:
144 <221> NAME/KEY: unsure

```

RAW SEQU LISTING D 06/24/2002
 PATENT APPLICATION: US/09/890,811A TIME: 17:40:08

Input Set : N:\Crf3\Refhold\I890811A.raw
 Output Set: N:\CRF3\06242002\I890811A.raw

145 <222> LOCATION: (640)
 146 <223> OTHER INFORMATION: n=a,c,g or t
 147 <220> FEATURE:
 148 <221> NAME/KEY: unsure
 149 <222> LOCATION: (653)
 150 <223> OTHER INFORMATION: n=a,c,g or t
 151 <220> FEATURE:
 152 <221> NAME/KEY: unsure
 153 <222> LOCATION: (680)
 154 <223> OTHER INFORMATION: n=a,c,g or t
 155 <220> FEATURE:
 156 <221> NAME/KEY: unsure
 157 <222> LOCATION: (692)
 158 <223> OTHER INFORMATION: n=a,c,g or t
 159 <220> FEATURE:
 160 <221> NAME/KEY: unsure
 161 <222> LOCATION: (703)
 162 <223> OTHER INFORMATION: n=a,c,g or t
 163 <400> SEQUENCE: 3
 164 gacgcacccc ntctntttctc tccccctctc gtctcgtecca gtcgtccccc tctccccccac 60
 165 tctaccacgc tgccctgctc tgcgtcgctc gccatggcgt cctccacggg gagcttggag 120
 166 cacggggggg tcaagtccac gccgcgcgcc ttcattacac ccttcaccga gctgctctcc 180
 167 ggcgcagggg acatgctagg agccggcgcc gatcaggagc ggtcgctgcc gagggggctg 240
 168 ttccaccgcg gcgccagggg cgtgccccaa ttcaagtcgg cgcagcctcc cagcctgccc 300
 169 atctgcgcgc cgcccattgc gccttcctcc tacttcgcca tcccgcccg gctcagcccc 360
 170 gccgagctgc tcgactcgcc cgtcctgctc cactcgtecc ccaacatcct ggcgctctcc 420
 171 accactggcg ccatcccggc gcagagggtc gactggaaga aggccgccga cctgatcgcg 480
 172 tctcagttct agcaagacgg cgacagccgg cgtgccgccc ccggttcga cgacttctcc 540
 173 ttcacacggg cactttcaac gccgtgcggc cgcacacgac gacgacgtcc ttaccttcat 600
 174 gaagaagaan gtggagaggg ccctggccga cgggcgcatt aacgcaaaat cnggtacaaa 660
 175 ggcgcggcac aaacaacccn aagccgggtg tncacgcgcc ggnaacttaa ttccgga 717
 177 <210> SEQ ID NO: 4
 178 <211> LENGTH: 150
 179 <212> TYPE: PRT
 180 <213> ORGANISM: Zea mays
 181 <400> SEQUENCE: 4
 182 Met Ala Ser Ser Thr Gly Ser Leu Glu His Gly Gly Phe Thr Phe Thr
 183 1 5 10 15
 184 Pro Pro Pro Phe Ile Thr Ser Phe Thr Glu Leu Leu Ser Gly Ala Gly
 185 20 25 30
 186 Asp Met Leu Gly Ala Gly Ala Asp Gln Glu Arg Ser Ser Pro Arg Gly
 187 35 40 45
 188 Leu Phe His Arg Gly Ala Arg Gly Val Pro Lys Phe Lys Ser Ala Gln
 189 50 55 60
 190 Pro Pro Ser Leu Pro Ile Ser Pro Pro Pro Met Ser Pro Ser Ser Tyr
 191 65 70 75 80
 192 Phe Ala Ile Pro Pro Gly Leu Ser Pro Ala Glu Leu Leu Asp Ser Pro
 193 85 90 95
 194 Val Leu Leu His Ser Ser Ser Asn Ile Leu Ala Ser Pro Thr Thr Gly

RAW SEQU LISTING D 06/24/2002
 PATENT APPLICATION: US/09/890,811A TIME: 17:40:08

Input Set : N:\Crf3\Refhold\I890811A.raw
 Output Set: N:\CRF3\06242002\I890811A.raw

```

195          100          105          110
196 Ala Ile Pro Ala Gln Arg Phe Asp Trp Lys Lys Ala Ala Asp Leu Ile
197          115          120          125
198 Ala Ser Gln Ser Gln Gln Asp Gly Asp Ser Arg Ala Ala Ala Ala Gly
199          130          135          140
200 Phe Asp Asp Phe Ser Phe
201          145          150
203 <210> SEQ ID NO: 5
204 <211> LENGTH: 1961
205 <212> TYPE: DNA
206 <213> ORGANISM: Oryza sativa
207 <400> SEQUENCE: 5
208 agtcgtctcgt ttctcgtctc cgatcactct cctcctcatc ttctgtcacgg tctcctcgtc 60
209 tcgctagctc gcttgcttgc tggctgagct gtggtacgct cgccatggcg tcctcgacgg 120
210 ggggggttga ccacgggttc acgttcacgc cgccgccgtt catcacgtcg ttcaccgagc 180
211 tgctgtcggg gggcgggtgg gacctgctcg gcgcggcgcg tgaggagcgc tcgccgaggg 240
212 ggttctccag aggcggagcg aggggtggcg gcgggggtgcc caagttcaag tcgcgcgacg 300
213 cgccgagcct gccgtctcgc ccgcgcggcg tgctgccgtc gtctacttc gccatcccgc 360
214 cggggctcag cccaccgag ctgctcgact ccccgtcct cctcagctcc tcccatatct 420
215 tggcggtccc gaccaccggt gcaatcccgc ctccagaggt cgactggaag gccagcgccg 480
216 atctcatcgc ttctcagcaa gatgacagcc gcggcgactt ctcttccac accaactccg 540
217 acgccaatgg cgcgcaaccg gcctctttcc ctctcttcaa ggagcaagag cagcaagtgg 600
218 tcgagtcgag caagaacggc gccgcgcgcg cgtcgagcaa caagagcggc ggcggcgagg 660
219 acaacaagct ggaggacggg tacaactgga ggaagtacgg gcagaagcag gtgaaggagg 720
220 gcgagaaccc gaggagctac tacaagtgca cctacaacgg ctgctccatg aagaagaagg 780
221 tggagcgctc gctcgcggac ggccgcacat cccagatcgt ctacaagggc gcacacaacc 840
222 accccaagcc gctctccacc gccgcaacgc ctcttccggc tccaccgccc ccgctgcgcg 900
223 cgacgacctc gcggcgccgc gcgcggcgcg ggaccagtac tccgcgcgca gccccgagaa 960
224 ctctctcgtc acgttcggcg acgacgaggg cgacaacgca tcgcaccgca gcgagggcga 1020
225 cgagcccga gccaagcgct ggaaaggagg atgctgacaa cgagggcgag tccggcgagg 1080
226 tgggcggcgg cgccggcggc aaccgggtgc gcgagccgag gcttggtgtg cagacgctga 1140
227 gcgacatcga catcctcgac aacggcttcc ggtggaggaa gtacggccag aaggtcgtca 1200
228 agggcaaccc caaccgaagg agctactaca agtgacgac ggtgggctgc ccggtgcgga 1260
229 agcacgtgga gcgggcgtcg cacgacacgc gcgccgtgat caccacctac gagggcaagc 1320
230 acaaccacga cgtcccggta cggccgcggc ggcggcgggc gacgcgcccc ggcgcggcgg 1380
231 tcgcctacgg ctgggggcga tccggggcga cggacgtcgc cgccgcccag caggggccct 1440
232 acaccctcga gatgctcccc aaccgcgcgc gcctctacgg cggctacggc gccggcgccg 1500
233 gcggcgccgc gttccgcgcg accaaggacg agcggcgagg cgacctgttc gtcgagtcgc 1560
234 tcctctgcta gtcgagccga gccgagccga gctgagctgg gccccacatc cccctgctcg 1620
235 ccacgtggcg tattttcgcc tcgcgtata cgtacggccg tatagcgtac gtatacacgc 1680
236 tcgcacgccc tgcccaacac ggcaatacac acatacatac tctcgtacac acgtagtagc 1740
237 atacatatac agtatagtag gtggtagtg tagctagcta gggagtgaga tccaatttgt 1800
238 tgattcggtt caggccactg ccacgtgggc cacaccggaa acagtacacg cgtatacacc 1860
239 aacttggga tacgcgtacg tacgcacatg tacacgtagt tttgtgcctt tgtaactgct 1920
240 gagagacagg tcaataaga ctgatgaatt tttcatttct t 1961
242 <210> SEQ ID NO: 6
243 <211> LENGTH: 488
244 <212> TYPE: PRT
245 <213> ORGANISM: Oryza sativa

```

RAW SEQUENCE LISTING ERROR SUMMARY D 06/24/2002
PATENT APPLICATION: US/09/890,811A TIME: 17:40:09

Input Set : N:\Crf3\Refhold\I890811A.raw
Output Set: N:\CRF3\06242002\I890811A.raw

Please Note:

Use of n and/or Xaa have been detected in the Sequence Listing. Please review the Sequence Listing to ensure that a corresponding explanation is presented in the <220> to <223> fields of each sequence which presents at least one n or Xaa.

Seq#:1; N Pos. 368,387,405,408,445,447,453,462,489,490,491,500
Seq#:2; Xaa Pos. 123,129,135,136,148,149,151,154,163,164,167
Seq#:3; N Pos. 11,15,610,640,653,680,692,703

Invalid Line Length:

The rules require that a line not exceed 72 characters in length. This includes spaces.

Seq#:14; Line(s) 743